

GenCore version 4.5
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OM protein - protein search, using sw model.

Run on: March 1, 2001, 15:46:57 ; Search time 210.42 Seconds
Sequence: 1 NQEDQTECQCQCRRCRQE..... RQQQCQRCKEICEEEERY 43
Scoring table: BLOSUM22
Gapop 10.0 , Gapext 0.5

Searched: 268485 seqs, 34193795 residues

Total number of hits satisfying chosen parameters: 268485

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : A_Geneseq_36:*

1: /SDS1/gcadata/geneseq/geneseq/AA1980.DAT:*

2: /SDS1/gcadata/geneseq/genescq/AA1981.DAT:*

3: /SDS1/gcadata/geneseq/geneseq/AA1982.DAT:*

4: /SDS1/gcadata/geneseq/genescq/AA1983.DAT:*

5: /SDS1/gcadata/geneseq/genescq/AA1984.DAT:*

6: /SDS1/gcadata/geneseq/genescq/AA1985.DAT:*

7: /SDS1/gcadata/geneseq/genescq/AA1986.DAT:*

8: /SDS1/gcadata/geneseq/genescq/AA1987.DAT:*

9: /SDS1/gcadata/geneseq/genescq/AA1988.DAT:*

10: /SDS1/gcadata/geneseq/genescq/AA1989.DAT:*

11: /SDS1/gcadata/geneseq/genescq/AA1990.DAT:*

12: /SDS1/gcadata/geneseq/genescq/AA1991.DAT:*

13: /SDS1/gcadata/geneseq/genescq/AA1992.DAT:*

14: /SDS1/gcadata/geneseq/genescq/AA1993.DAT:*

15: /SDS1/gcadata/geneseq/genescq/AA1994.DAT:*

16: /SDS1/gcadata/geneseq/genescq/AA1995.DAT:*

17: /SDS1/gcadata/geneseq/genescq/AA1996.DAT:*

18: /SDS1/gcadata/geneseq/genescq/AA1997.DAT:*

19: /SDS1/gcadata/geneseq/genescq/AA1998.DAT:*

20: /SDS1/gcadata/geneseq/genescq/AA1999.DAT:*

21: /SDS1/gcadata/geneseq/genescq/AA2000.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match Length	DB ID	Description
1	248	100.0	666 19	W62828
2	242	97.6	666 19	W62829
3	241	97.2	625 19	W62830
4	117	47.2	525 19	W62831
5	117	47.2	566 13	R20181
6	109	44.0	590 19	W62832
7	89	35.9	28 19	W62841
8	66.5	26.8	593 19	W62835
9	66.5	26.8	637 19	W62837
10	65	26.2	919 10	P93109
11	65	26.2	919 18	W11783
12	65	26.2	919 21	Y78914

13: 64.5 26.0 35 13 R21079 Antimicrobial maize
14: 63.5 25.6 33 19 W62836 Zea mays antinicro
15: 63.5 25.6 51 18 W33694 Mouse protamine 1.
16: 63.5 25.6 176 18 W33695 Mouse protamine 1.
17: 63.5 25.6 301 19 W37085 Anti-human SC sing
18: 63 25.4 910 20 Y22191 Mouse brain CNG-1
19: 62.5 25.2 107 17 R91705
20: 62.5 25.2 107 20 Y30404
21: 62.5 25.2 342 20 Y16785
22: 60.5 24.4 215 14 R44806
23: 60 24.2 154 20 Y33504
24: 60 24.2 918 12 R12223
25: 60 24.2 918 20 Y33491
26: 59.5 24.0 98 21 Y65429 Human androgen rec
27: 59.5 23.8 445 21 Y32874 Human secreted pro
28: 59 23.8 147 17 W03226 Human cyclin D3 ps
29: 59 23.8 147 20 W81029 Murine PCIP prote
30: 58.5 23.6 88 20 P93116 Human unliganded a
31: 58.5 23.6 514 19 W80400 Human androgen rec
32: 58 23.4 151 21 Y74634 Human 5'-EST relat
33: 58 23.2 199 21 Y74635 Neisseria meningit
34: 57.5 23.2 304 13 R20063 LXR-alpha, orphan
35: 57.5 23.2 317 10 P93116 Murine PCIP prote
36: 57.5 23.2 809 20 Y29672 Mature nematode ex
37: 57 23.0 303 15 R60054 Dirofilaria immiti
38: 57 23.0 440 14 R33744 XR2. Homo sapiens
39: 57 23.0 567 20 Y22212 MTG16b protooncogene
40: 57 23.0 653 20 Y22211 MTG16a protein seq
41: 57 23.0 704 20 Y22209 AMLI-MTG16 fusion
42: 57 23.0 752 15 R51701 AMLI-MTG16 fusion
43: 57 23.0 780 20 Y22210 Amino acid sequenc
44: 57 23.0 780 21 Y54320 Human cytoskeleton
45: 56.5 22.8 281 21 Y91958

ALIGNMENTS

RESULT	1.
ID	W62828 standard; protein; 666 AA.
XX	W62828;
AC	XX
DT	27-OCT-1998 (first entry)
DE	Macadamia integrifolia antimicrobial protein.
KW	antimicrobial protein; infestation; control.
OS	Macadamia integrifolia.
XX	
Key	FII Location/Qualifiers
FT	Peptide 1..28
FT	/note= "signal peptide"
FT	29..666
FT	/note= "mature protein"
PN	W09827805-A1.
XX	
PD	02-JUL-1998.
XX	
PF	22-DEC-1997; 97NO-AU00874.
PR	20-DEC-1996; 96AU-0004275.
XX	
PA	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX	
PI	Bower NI, Goultier KC, Green JL, Manners JM, Marcus JP;
XX	
DR	WP1; 1998-37279/32.
DR	N-PSDB; V42310.
XX	

Human androgen rec
Nematode extracted
Mouse protamine 1.
Mouse protamine 1.
Anti-human SC sing
Mouse brain CNG-1
Acanap23. Acanlos
Human secreted pro
Human cyclin D3 ps
Human unliganded a
Human androgen rec
Human androgen rec
Human 5'-EST relat
Neisseria meningit
LXR-alpha, orphan
Murine PCIP prote
Mature nematode ex
A secreted protein
Neisseria meningit
XR2. Homo sapiens
MTG16b protooncogene
MTG16a protein seq
AMLI-MTG16 fusion
AMLI-MTG16 fusion
Amino acid sequenc
Human cytoskeleton

PT Novel anti-microbial protein from e.g. Macadamia integrifolia -
 PT useful for controlling microbial infestations of plants or mammals.
 XX
 PS Claim 1; Page 34-36; 96pp; English.
 XX
 CC The sequence is that of an antimicrobial protein which can
 CC be used to control microbial infestations in plants and mammalian
 CC animals.
 XX
 SQ Sequence 666 AA:

Query Match 100.0%; Score 248; DB 19; Length 666;
 Best Local Similarity 100.0%; Pred. No. 5 8e-20; Mismatches 0; Indels 0; Gaps 0;
 Matches 43; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 NQEDPOTECQQCQRRCQEQESPRQQYCQRCKECEEEEEEY 43
 Db 74 nqedpqteccqgqcrqrcrqrqesprqqycqrckelceeeey 116

RESULT 2
 ID W62829 standard; Protein; 666 AA.
 AC W62829;
 XX
 DT 27-OCT-1998 (first entry)
 DE Macadamia integrifolia antimicrobial protein.
 KW antimicrobial protein; infestation; control.
 OS Macadamia integrifolia.
 XX
 FH Key Location/Qualifiers
 FT Peptide 1..28
 FT Protein /note= "signal peptide"
 FT /note= "mature protein"
 XX
 PN WO9827805-A1.
 XX
 PD 02-JUL-1998.
 XX
 PR 22-DEC-1997; 97WO-AU00874.
 XX
 PR 20-DEC-1996; 96AU-0004275.
 XX
 PA (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
 XX
 PI Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
 XX
 DR WO1998377279/32.
 XX
 DR N-PSDB; V42316.
 XX
 PS Novel anti-microbial protein from e.g. Macadamia integrifolia -
 PT useful for controlling microbial infestations of plants or mammals
 XX
 PS claim 1; Page 43-45; 96pp; English.
 XX
 CC The sequence is that of an antimicrobial protein which can
 CC be used to control microbial infestations in plants and mammalian
 CC animals.
 XX
 SQ Sequence 625 AA:

Query Match 97.2%; Score 241; DB 19; Length 625;
 Best Local Similarity 97.7%; Pred. No. 3 2e-19; Mismatches 1; Indels 0; Gaps 0;
 Matches 42; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 NQEDPOTECQQCQRRCQEQESPRQQYCQRCKECEEEEEEY 43
 Db 33 nqedpqteccqgqcrqrcrqrqesprqqycqrckelceeeey 75

RESULT 4
 ID W62831 standard; Protein; 525 AA.
 AC W62831;
 XX
 DT 27-OCT-1998 (first entry)
 DE Theobroma cacao antimicrobial protein.
 KW antimicrobial protein; infestation; control.
 XX
 OS Theobroma cacao.

Query Match 97.6%; Score 242; DB 19; Length 666;
 Best Local Similarity 95.3%; Pred. No. 2.7e-19;
 Matches 41; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 1 NQEDPOTECQQCQRRCQEQESPRQQYCQRCKECEEEEEEY 43

XX	WO9827805-A1.	CC	derived from the 67 kD precursor. T:
XX	PN	detected in a cDNA library prepared from immature cocoa beans RNA	
XX	PD	using a probe based on the AA sequence of a CNBR peptide common to	
XX	PR	the 47 kD and 31 kD polypeptides. Homology searches revealed close	
XX	PF	homologies between the 67 kD polypeptide and the vicilins, which are	
XX	22-DEC-1997;	CC	seed storage proteins.
XX	20-DEC-1996;	CC	Sequence 566 AA;
XX	PR	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.	
XX	PA	Query Match Similarity 47.2%; Score 117; DB 13; Length 566;	
XX	PI	Best Local Similarity 50.0%; Pred. No. 1 3e-05; Indels 0; Gaps 0;	
XX	DR	Matches 20; Conservative 10; Mismatches 10; Indels 0; Gaps 0;	
XX	WPI;	Claim 1; Page 47-49; 96pp; English.	
XX	1998-377279/32.	The sequence is that of an antimicrobial protein which can	
CC	be used to control microbial infestations in plants and mammalian		
CC	animals.		
XX	CC	Novel anti-microbial protein from e.g. Macadamia integrifolia -	
XX	PT	useful for controlling microbial infestations of plants or mammals	
XX	PR	Sequence 525 AA;	
XX	PS	Query Match Similarity 47.2%; Score 117; DB 13; Length 525;	
XX	PI	Best Local Similarity 50.0%; Pred. No. 1 2e-05; Indels 0; Gaps 0;	
XX	DR	Matches 20; Conservative 10; Mismatches 10; Indels 0; Gaps 0;	
XX	WPI;	Claim 1; Page 47-49; 96pp; English.	
XX	1998-377279/32.	The sequence is that of an antimicrobial protein which can	
CC	be used to control microbial infestations in plants and mammalian		
CC	animals.		
XX	CC	Novel anti-microbial protein from e.g. Macadamia integrifolia -	
XX	PT	useful for controlling microbial infestations of plants or mammals	
XX	PR	Sequence 525 AA;	
XX	PS	Query Match Similarity 47.2%; Score 117; DB 13; Length 566;	
XX	PI	Best Local Similarity 50.0%; Pred. No. 1 3e-05; Indels 0; Gaps 0;	
XX	DR	Matches 20; Conservative 10; Mismatches 10; Indels 0; Gaps 0;	
XX	WPI;	Claim 1; Page 47-49; 96pp; English.	
XX	1998-377279/32.	The sequence is that of an antimicrobial protein which can	
CC	be used to control microbial infestations in plants and mammalian		
CC	animals.		
XX	CC	Novel anti-microbial protein from e.g. Macadamia integrifolia -	
XX	PT	useful for controlling microbial infestations of plants or mammals	
XX	PR	Sequence 590 AA;	
XX	PS	Query Match Similarity 44.0%; Score 109; DB 19; Length 590;	
XX	PI	Best Local Similarity 50.0%; Pred. No. 0.0001; Indels 2; Gaps 1;	
XX	DR	Matches 20; Conservative 8; Mismatches 10; Indels 2; Gaps 1;	
XX	WPI;	Claim 4; Fig 2; 59pp; English.	
XX	N-PSDB; Q20377.	The inventors claim a 67 kD and 31 kD T. cacao protein, and	
CC	Recombinant cacao proteins - are responsible for flavour in cacao		
PT	beans and produced in large quantities using yeast and bacterial		
PT	expression vectors		
PT	Spencer ME, Hodge R, Peakin EA, Ashton S;		
PT	WPI: 1992-024118/03.		
PT	N-PSDB; Q20377.		
XX	3 EDPOTECOCQCQRROQESGPROQQYCQRCKELCEEEEEE 42		
XX	Db 81 edparryyecqccqrqee-rgdpqccqrkclifeqeqq 118		
XX	RESULT 7		
XX	W62841		
XX	ID W62841 standard; Protein: 28 AA.		
XX	AC W62841;		
AC	W62841;		

CC	androgen-independent activation of the AR. The androgen receptor acts as a transcription factor, regulating the expression of certain androgen-responsive genes. Interaction of the AR with the protein kinase A signal transduction pathway involves interaction with the androgen independent region. The AR fragment and peptides derived from it can be used as agents for inhibiting androgen independent activation of the androgen receptor, as activation domains, and as a tool for screening for compounds which affect androgen independent activation of the AR. The peptides, when used in combination with androgen deprivation, effectively limit androgen mediated disease progression. These diseases include cancer, benign prostatic hyperplasia, hirsutism, androgenic alopecia, acne, breast cancer, Kennedy disease, and especially prostate cancer. The peptides and nucleic acids encoding them, are especially used for the treatment of androgen-mediated diseases, especially prostate tumours in patients deprived of androgen.
CC	Sequence 919 AA;
CC	SQ
CC	Query Match 26.2%; Score 65; DB 21; Length 919;
CC	Best Local Similarity 48.3%; Pred. No. 11; Mismatches 9; Indels 0; Gaps 0;
CC	Matches 14; Conservative 6; Mismatches 9; Indels 0; Gaps 0;
OY	2 QEDPOTECQCQRRCRQESPRQQYCQ 30
OY	1: - : : ; : -
Db	61 .qqqqqqqqqqqqqqqqqgetsprqqqqq 89
RESULT	13
R21079	
ID	R21079 standard; Peptide; 35 AA.
XX	
XX	R21079;
AC	
XX	DPT 09-APR-1992 (first entry)
XX	DE Antimicrobial maize peptide, CMIII.
XX	Maize; CMIII; corn; pathogen.
KW	
XX	OS zea mays.
OS	
XX	XX
PN	EP465009-A.
XX	
PD	08-JAN-1992.
XX	
PF	05-JUN-1991; 91EP-0305054.
XX	
PR	05-JUN-1990; 90US-0536127.
XX	
PA	(PION-) PIONEER HI-BRED INT.
XX	
PT	Duvick JP, Rood TA, Rao AG;
XX	
DR	WPI; 1992-010214/02.
XX	
PT	Use of maize seed peptide CMIII and DNA encoding it - for killing or inhibiting plant pathogenic microorganisms.
PT	Example 2; Page 5; Zipp; English.
PS	
XX	The peptide (SEQ ID NO 1) was purified from public corn variety B73 and proprietary corn variety MH8. It is basic and has a total mol. wt. of 3900 daltons. The peptide sequence was used to design probes which were used to screen a maize genomic or cDNA library. The isolated CMIII gene can be used to prepare an expression vector for produc. of recombinant CMIII for use in controlling plant patho- genic organisms. See also Q20272 and 3.
CC	
CC	See also Q20272 and 3.
CC	
CC	Sequence 35 AA;
XX	

Query Match		Score 64.5;	DB 13;	Length 35;
Best Local Similarity		44.0%	Pred. No. 0.57;	
Matches		11;	Conservative	Mismatches
DE		6		
RESULT	14			
W62836				
ID	W62836	standard; Protein:	33 AA.	
XX				
AC	W62836;			
XX				
DT	27-OCT-1998	(first entry)		
XX				
DE	zea mays antimicrobial protein.			
XX				
KW	antimicrobial protein; infestation; control.			
XX				
OS	zea mays.			
XX				
PN	W09827805-A1.			
XX				
PD	02-JUL-1998.			
XX				
PF	22-DBC-1997;	97WO-AU00874.		
XX				
PR	20-DEC-1996;	96AU-0004275.		
XX				
PA	(RETR-) COOP RBS CENT TROPICAL PLANT PATHOLOGY.			
XX				
PI	Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;			
XX				
DR	WPI: 1998-377279/32.			
XX				
PT	Novel anti-microbial protein from e.g. Macadamia integrifolia - useful for controlling microbial infestations of plants or mammals			
XX				
PS	Disclosure; Page 60; 96pp; English.			
XX				
CC	The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian animals.			
CC				
XX				
SQ	Sequence 33 AA;			
Query Match		25.6%; Score 63.5;	DB 19;	Length 33;
Best Local Similarity		44.0%; Pred. No. 0.7;		
Matches		6; Mismatches	7;	Indels
DE		1;	Gaps	1;
OY	11 QCQRC-RQQESGRQQYQQRCK 34			
Db	: : : : : :			
	6 ecrrrqclrrheggqpwtqcemrcc 30			
RESULT	15			
W33694				
ID	W33694 standard; peptide;	51 AA.		
XX				
AC	W33694;			
XX				
DT	30-APR-1998 (first entry)			
XX				
DE	Mouse protamine 1.			
XX				
KW	Nucleic acid-binding motif; HBC; protamine 1; mouse; diagnosis;			
KW	HCV core polypeptide; immunoassay; detection; antigen; disease;			
KW	Hepatitis C virus; infection.			
XX				

OS
XX
PN
XX
EP805160-A1.
PD
XX
05-NOV-1997.
PF
XX
30-APR-1997; 97EP-0400985.
PR
XX
01-MAY-1996; 96JP-0134444.
PA
(FURE) FUJIREBIO INC.
XX
PI
Itoh S, Takemura F, Ueno E;
XX
WPI; 1997-529030/49.
DR
DR
N-PSDB; v06342.
XX
PT
Nucleic acid-bound polypeptide - useful as immunoassay reagent.
XX
PS
Example 15; Pages 32-33; 38pp; English.
XX
CC
This is a mouse protamine 1. This is used in a method for producing a fusion
CC nucleic acid-bound polypeptide. The method comprises producing a fusion
gene containing the polypeptide and a nucleic acid-binding motif, binding
CC a nucleic acid to the polypeptide as a soluble fraction, and purifying the
CC nucleic acid-bound polypeptide from the soluble fraction. When the
CC polypeptide is a recombinant form of an antigen, the nucleic acid-bound
CC antigen or an antibody to the antigen, especially in an agglutination
CC assay using particles coated with the nucleic acid-bound polypeptide. The
CC methods can be applied to diagnosis of disease and infection, especially
CC for the detection of HBV and HCV polypeptides. The nucleic acid-bound
CC polypeptides may be immunoreactive in cases where the free polypeptide
CC is not.
XX
Sequence 51 AA;
SQ

Query Match 25.6%; Score 63.5; DB 18; Length 51;
Best Local Similarity 28.9%; Pred. No. 1;
Matches 11; Conservative 11; Mismatches 11; Indels 5; Gaps 1;
OY 6 QTCGQOCORRCRQQESGPROQQCQRREKEICPEEEY 43
Db :: |:: |||: |:: ||| : | | |
12 rsrccrrrccrr---rrccrrrccrrrrsy 44

Search completed: March 1, 2001, 15:46:59
Job time: 224 sec

